B1

SEQUENCE LISTING

<110> Danks, Mary K.
Potter, Philip M.
Houghton, Peter J.

<120> Compositions and Methods for Sensitizing and Inhibiting Growth of Human Tumor Cells

<130> SJ-0011

<140> 09/622,568
<141> 2000-08-31

<150> 60/075,258
<151> 1998-02-19

<150> PCT/US99/03171
<151> 1999-02-12

<160> 30

<170> PatentIn Ver. 2.0

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Gly Lys Phe Val Ser Xaa Glu Gly Phe Ala Gln Pro Val Ala Lys Phe
20 25 30

Xaa Gly

<210> 2
<211> 36
<212> PRT
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 2
His Pro Ser Ala Pro Pro Val Val Asp Thr Val Lys Gly Lys Val Leu
1 5 10 15

Gly Lys Phe Val Ser Leu Glu Gly Phe Ala Gln Pro Val Ala Val Phe
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Leu Gly Val Pro
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<210> 3
<211> 54
<212> PRT
<213> Homo sapiens

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 3
Met Trp Leu Arg Ala Phe Ile Leu Ala Thr Leu Ser Ala Ser Ala Ala
1 5 10 15

Trp Gly His Pro Ser Ser Pro Pro Val Val Asp Thr Val His Gly Lys
20 25 30

Val Leu Gly Lys Phe Val Ser Leu Glu Gly Phe Ala Gln Pro Val Ala
35 40 45

Ile Phe Leu Gly Ile Pro
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<210> 4
<211> 54
<212> PRT
<213> Rattus rattus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 4
Met Trp Leu Cys Ala Leu Val Trp Ala Ser Leu Ala Val Cys Pro Ile
1 5 10 15

Trp Gly His Pro Ser Ser Pro Pro Val Val Asp Thr Thr Lys Gly Lys
20 25 30

Val Leu Gly Lys Tyr Val Ser Leu Glu Gly Phe Thr Gln Pro Val Ala
35 40 45

Val Phe Leu Gly Val Pro
50

<210> 5
<211> 54
<212> PRT
<213> Mus musculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 5
Met Trp Leu His Ala Leu Val Trp Ala Ser Leu Ala Val Cys Pro Ile
1 5 10 15

Leu Gly His Ser Leu Leu Pro Pro Val Val Asp Thr Thr Gln Gly Lys
20 25 30

Val Leu Gly Lys Tyr Ile Ser Leu Glu Gly Phe Glu Gln Pro Val Ala
35 40 45

Val Phe Leu Gly Val Pro
50

<210> 6
<211> 5
<212> PRT
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 6
His Pro Ser Ala Pro
1 5

<210> 7
<211> 14
<212> DNA
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 7
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<210> 8
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
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<400> 8
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<210> 9
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
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<400> 9
cacccntcng cncc 14

<210> 10
<211> 7
<212> PRT
<213> Oryctolagus cuniculus

<220>
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<400> 10
Ala Phe Trp Thr Glu Leu Trp
1 5

<210> 11
<211> 21
<212> DNA
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 11
gcattctgga cagaactatg g 21

<210> 12
<211> 21
<212> DNA
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 12
ccaaagttca gtccagaaag c 21

<210> 13
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
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<400> 13
ccanagttcn gtccagaang c 21

<210> 14
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
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<223> i

<400> 14
ccataattcn gtccagaang c 21

<210> 15
<211> 30
<212> PRT
<213> Rattus rattus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 15
Met Trp Leu Cys Ala Leu Ala Leu Ala Ser Leu Ala Ala Cys Thr Ala
1 5 10 15

Trp Gly His Pro Ser Ala Pro Pro Val Val Asp Thr Val Lys
20 25 30

<210> 16
<211> 30
<212> PRT
<213> Rattus sp.

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 16
Met Trp Leu Cys Ala Leu Val Trp Ala Ser Leu Ala Val Cys Pro Ile
1 5 10 15

Trp Gly His Pro Ser Ser Pro Pro Val Val Asp Thr Thr Lys
20 25 30

<210> 17
<211> 30
<212> PRT
<213> Homo sapiens

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 17
Met Trp Leu Arg Ala Phe Ile Leu Ala Thr Leu Ser Ala Ser Ala Ala
1 5 10 15

Trp Gly His Pro Ser Ser Pro Pro Val Val Asp Thr Val His
20 25 30

<210> 18
<211> 30
<212> PRT
<213> Rattus rattus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 18
Met Arg Leu Tyr Pro Leu Val Trp Leu Phe Leu Ala Ala Cys Thr Ala
1 5 10 15

Trp Gly Tyr Pro Ser Ser Pro Pro Val Val Asn Thr Val Lys
20 25 30

<210> 19
<211> 30
<212> PRT
<213> Mus musculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 19
Met Trp Leu His Ala Leu Val Trp Ala Ser Leu Ala Val Cys Pro Ile
1 5 10 15

Leu Gly His Ser Leu Leu Pro Pro Val Val Asp Thr Thr Gln
20 25 30

<210> 20
<211> 1717
<212> DNA
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 20
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ggggcacccg tctgcaccac ctgtggtaga tactgtgcat ggcaaagtcc tggggaagtt 120

cgtcagctta gaaggatttg cacagcccgt ggccgtcttc ctgggagtcc ccttcgccaa 180
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gaagaacacc acctcctacc ctcccatgtg ctcccaggac gcagtatcag ggcatatgct 300
ctcggagctc ttcaccaaca gaaaagagaa catccctctt aagttttctg aagactgcct 360
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catgagaccc aagacagtga taggggacca tggagatgag atcttctctg tcttaggagc 1440
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actgaaagac aaggaagtgg ctttctggac tgagctctgg gccaaggagg cagcaaggcc 1680
acgtgagaca gagcacattg agctgtgaat tgaattc 1717

<210> 21
<211> 565
<212> PRT
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 21
Met Trp Leu Cys Ala Leu Ala Leu Ala Ser Leu Ala Ala Cys Thr Ala
1 5 10 15

Trp Gly His Pro Ser Ala Pro Pro Val Val Asp Thr Val His Gly Lys
20 25 30

Val Leu Gly Lys Phe Val Ser Leu Glu Gly Phe Ala Gln Pro Val Ala
35 40 45

Val Phe Leu Gly Val Pro Phe Ala Lys Pro Pro Leu Gly Ser Leu Arg
50 55 60

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Phe Ala Pro Pro Gln Pro Ala Glu Ser Trp Ser His Val Lys Asn Thr
 65                  70                  75                  80

Thr Ser Tyr Pro Pro Met Cys Ser Gln Asp Ala Val Ser Gly His Met
                 85                  90                  95

Leu Ser Glu Leu Phe Thr Asn Arg Lys Glu Asn Ile Pro Leu Lys Phe
            100                 105                 110

Ser Glu Asp Cys Leu Tyr Leu Asn Ile Tyr Thr Pro Ala Asp Leu Thr
        115                 120                 125

Lys Arg Gly Arg Leu Pro Val Met Val Trp Ile His Gly Gly Gly Leu
    130                 135                 140

Met Val Gly Gly Ala Ser Thr Tyr Asp Gly Leu Ala Leu Ser Ala His
145                 150                 155                 160

Glu Asn Val Val Val Val Thr Ile Gln Tyr Arg Leu Gly Ile Trp Gly
                165                 170                 175

Phe Phe Ser Thr Gly Asp Glu His Ser Arg Gly Asn Trp Gly His Leu
            180                 185                 190

Asp Gln Val Ala Ala Leu Arg Trp Val Gln Asp Asn Ile Ala Asn Phe
        195                 200                 205

Gly Gly Asp Pro Gly Ser Val Thr Ile Phe Gly Glu Ser Ala Gly Gly
    210                 215                 220

Gln Ser Val Ser Ile Leu Leu Leu Ser Pro Leu Thr Lys Asn Leu Phe
225                 230                 235                 240

His Arg Ala Ile Ser Glu Ser Gly Val Ala Leu Leu Ser Ser Leu Phe
                245                 250                 255

Arg Lys Asn Thr Lys Ser Leu Ala Glu Lys Ile Ala Ile Glu Ala Gly
            260                 265                 270

Cys Lys Thr Thr Thr Ser Ala Val Met Val His Cys Leu Arg Gln Lys
        275                 280                 285

Thr Glu Glu Glu Leu Met Glu Val Thr Leu Lys Met Lys Phe Met Ala
    290                 295                 300

Leu Asp Leu Val Gly Asp Pro Lys Glu Asn Thr Ala Phe Leu Thr Thr
305                 310                 315                 320
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Val Ile Asp Gly Val Leu Leu Pro Lys Ala Pro Ala Glu Ile Leu Ala
                325                 330                 335

Glu Lys Lys Tyr Asn Met Leu Pro Tyr Met Val Gly Ile Asn Gln Gln
            340                 345                 350

Glu Phe Gly Trp Ile Ile Pro Met Gln Met Leu Gly Tyr Pro Leu Ser
        355                 360                 365

Glu Gly Lys Leu Asp Gln Lys Thr Ala Thr Glu Leu Leu Trp Lys Ser
    370                 375                 380

Tyr Pro Ile Val Asn Val Ser Lys Glu Leu Thr Pro Val Ala Thr Glu
385                 390                 395                 400

Lys Tyr Leu Gly Gly Thr Asp Asp Pro Val Lys Lys Lys Asp Leu Phe
                405                 410                 415

Leu Asp Met Leu Ala Asp Leu Leu Phe Gly Val Pro Ser Val Asn Val
            420                 425                 430

Ala Arg His His Arg Asp Ala Gly Ala Pro Thr Tyr Met Tyr Glu Tyr
        435                 440                 445

Arg Tyr Arg Pro Ser Phe Ser Ser Asp Met Arg Pro Lys Thr Val Ile
    450                 455                 460

Gly Asp His Gly Asp Glu Ile Phe Ser Val Leu Gly Ala Pro Phe Leu
465                 470                 475                 480

Lys Glu Gly Ala Thr Glu Glu Glu Ile Lys Leu Ser Lys Met Val Met
                485                 490                 495

Lys Tyr Trp Ala Asn Phe Ala Arg Asn Gly Asn Pro Asn Gly Glu Gly
            500                 505                 510

Leu Pro Gln Trp Pro Ala Tyr Asp Tyr Lys Glu Gly Tyr Leu Gln Ile
        515                 520                 525

Gly Ala Thr Thr Gln Ala Ala Gln Lys Leu Lys Asp Lys Glu Val Ala
    530                 535                 540

Phe Trp Thr Glu Leu Trp Ala Lys Glu Ala Ala Arg Pro Arg Glu Thr
545                 550                 555                 560

Glu His Ile Glu Leu
                565
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<210> 22
<211> 6
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 22
cacgtg 6

<210> 23
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 23
ggcaggaatt ctgccatgtg gctctg 26

<210> 24
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 24
cgggaattca cattcacagc tcaatgt 27

<210> 25
<211> 6
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 25
cacctg 6

<210> 26
<211> 543

<212> PRT
<213> Oryctolagus cuniculus

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 26
Met Trp Leu Cys Ala Leu Ala Leu Ala Ser Leu Ala Ala Cys Thr Ala
1 5 10 15

Trp Gly His Pro Ser Ala Pro Pro Val Val Asp Thr Val His Gly Lys
20 25 30

Val Leu Gly Lys Phe Val Ser Leu Glu Gly Phe Ala Gln Pro Val Ala
35 40 45

Val Phe Leu Gly Val Pro Phe Ala Lys Pro Pro Leu Gly Ser Leu Arg
50 55 60

Phe Ala Pro Pro Gln Pro Ala Glu Ser Trp Ser His Val Lys Asn Thr
65 70 75 80

Thr Ser Tyr Pro Pro Met Cys Ser Gln Asp Ala Val Ser Gly His Met
85 90 95

Leu Ser Glu Leu Phe Thr Asn Arg Lys Glu Asn Ile Pro Leu Lys Phe
100 105 110

Ser Glu Asp Cys Leu Tyr Leu Asn Ile Tyr Thr Pro Ala Asp Leu Thr
115 120 125

Lys Arg Gly Arg Leu Pro Val Met Val Trp Ile His Gly Gly Gly Leu
130 135 140

Met Val Gly Gly Ala Ser Thr Tyr Asp Gly Leu Ala Leu Ser Ala His
145 150 155 160

Glu Asn Val Val Val Val Thr Ile Gln Tyr Arg Leu Gly Ile Trp Gly
165 170 175

Phe Phe Ser Thr Gly Asp Glu His Ser Arg Gly Asn Trp Gly His Leu
180 185 190

Asp Gln Val Ala Ala Leu Arg Trp Val Gln Asp Asn Ile Ala Asn Phe
195 200 205

Gly Gly Asp Pro Gly Ser Val Thr Ile Phe Gly Glu Ser Ala Gly Gly
210 215 220

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Gln Ser Val Ser Ile Leu Leu Leu Ser Pro Leu Thr Lys Asn Leu Phe
225                 230                 235                 240

His Arg Ala Ile Ser Glu Ser Gly Val Ala Leu Leu Ser Ser Leu Phe
                245                 250                 255

Arg Lys Asn Thr Lys Ser Leu Ala Glu Lys Ile Ala Ile Glu Ala Gly
            260                 265                 270

Cys Lys Thr Thr Thr Ser Ala Val Met Val His Cys Leu Arg Gln Lys
        275                 280                 285

Thr Glu Glu Glu Leu Met Glu Val Thr Leu Lys Met Lys Phe Met Ala
    290                 295                 300

Leu Asp Leu Val Gly Asp Pro Lys Glu Asn Thr Ala Phe Leu Thr Thr
305                 310                 315                 320

Val Ile Asp Gly Val Leu Leu Pro Lys Ala Pro Ala Glu Ile Leu Ala
                325                 330                 335

Glu Lys Lys Tyr Asn Met Leu Pro Tyr Met Val Gly Ile Asn Gln Gln
            340                 345                 350

Glu Phe Gly Trp Ile Ile Pro Met Gln Met Leu Gly Tyr Pro Leu Ser
        355                 360                 365

Glu Gly Lys Leu Asp Gln Lys Thr Ala Thr Glu Leu Leu Trp Lys Ser
    370                 375                 380

Tyr Pro Ile Val Asn Val Ser Lys Glu Leu Thr Pro Val Ala Thr Glu
385                 390                 395                 400

Lys Tyr Leu Gly Gly Thr Asp Asp Pro Val Lys Lys Lys Asp Leu Phe
                405                 410                 415

Leu Asp Met Leu Ala Asp Leu Leu Phe Gly Val Pro Ser Val Asn Val
            420                 425                 430

Ala Arg His His Arg Asp Ala Gly Ala Pro Thr Tyr Met Tyr Glu Tyr
        435                 440                 445

Arg Tyr Arg Pro Ser Phe Ser Ser Asp Met Arg Pro Lys Thr Val Ile
    450                 455                 460

Gly Asp His Gly Asp Glu Ile Phe Ser Val Leu Gly Ala Pro Phe Leu
465                 470                 475                 480
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Lys Glu Gly Ala Thr Glu Glu Glu Ile Lys Leu Ser Lys Met Val Met
485 490 495

Lys Tyr Trp Ala Asn Phe Ala Arg Asn Gly Asn Pro Asn Gly Glu Gly
500 505 510

Leu Pro Gln Trp Pro Ala Tyr Asp Tyr Lys Glu Gly Tyr Leu Gln Ile
515 520 525

Gly Ala Thr Thr Gln Ala Ala Gln Lys Leu Lys Asp Lys Glu Val
530 535 540

<210> 27
<211> 2191
<212> DNA
<213> Homo sapiens

<220>
<223> Description of Artificial Sequence: Synthetic

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atccagtacc gcctgggtgt cctgggcttc ttcagcactg gagacaagca cgcaaccggc 660
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<210> 28
<211> 559
<212> PRT
<213> Homo sapiens

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 28
Met Arg Leu His Arg Leu Arg Ala Arg Leu Ser Ala Val Ala Cys Gly
1 5 10 15

Leu Leu Leu Leu Leu Val Arg Gly Gln Gly Gln Asp Ser Ala Ser Pro
20 25 30

Ile Arg Thr Thr His Thr Gly Gln Val Leu Gly Ser Leu Val His Val
35 40 45

Lys Gly Ala Asn Ala Gly Val Gln Thr Phe Leu Gly Ile Pro Phe Ala
50 55 60

Lys Pro Pro Leu Gly Pro Leu Arg Phe Ala Pro Pro Glu Pro Pro Glu
65 70 75 80

Ser Trp Ser Gly Val Arg Asp Gly Thr Thr His Pro Ala Met Cys Leu
85 90 95

Gln Asp Leu Thr Ala Val Glu Ser Glu Phe Leu Ser Gln Phe Asn Met
100 105 110

Thr Phe Pro Ser Asp Ser Met Ser Glu Asp Cys Leu Tyr Leu Ser Ile
115 120 125

Tyr Thr Pro Ala His Ser His Glu Gly Ser Asn Leu Pro Val Met Val
130 135 140

```
Trp Ile His Gly Gly Ala Leu Val Phe Gly Met Ala Ser Leu Tyr Asp
145                 150                 155                 160

Gly Ser Met Leu Ala Ala Leu Glu Asn Val Val Val Val Ile Ile Gln
                165                 170                 175

Tyr Arg Leu Gly Val Leu Gly Phe Phe Ser Thr Gly Asp Lys His Ala
            180                 185                 190

Thr Gly Asn Trp Gly Tyr Leu Asp Gln Val Ala Ala Leu Arg Trp Val
        195                 200                 205

Gln Gln Asn Ile Ala His Phe Gly Gly Asn Pro Asp Arg Val Thr Ile
    210                 215                 220

Phe Gly Glu Ser Ala Gly Gly Thr Ser Val Ser Ser Leu Val Val Ser
225                 230                 235                 240

Pro Ile Ser Gln Gly Leu Phe His Gly Ala Ile Met Glu Ser Gly Val
                245                 250                 255

Ala Leu Leu Pro Gly Leu Ile Ala Ser Ser Ala Asp Val Ile Ser Thr
            260                 265                 270

Val Val Ala Asn Leu Ser Ala Cys Asp Gln Val Asp Ser Glu Ala Leu
        275                 280                 285

Val Gly Cys Leu Arg Gly Lys Ser Lys Glu Glu Ile Leu Ala Ile Asn
    290                 295                 300

Lys Pro Phe Lys Met Ile Pro Gly Val Val Asp Gly Val Phe Leu Pro
305                 310                 315                 320

Arg His Pro Gln Glu Leu Leu Ala Ser Ala Asp Phe Gln Pro Val Pro
                325                 330                 335

Ser Ile Val Gly Val Asn Asn Asn Glu Phe Gly Trp Leu Ile Pro Lys
            340                 345                 350

Val Met Arg Ile Tyr Asp Thr Gln Lys Glu Met Asp Arg Glu Ala Ser
        355                 360                 365

Gln Ala Ala Leu Gln Lys Met Leu Thr Leu Leu Met Leu Pro Pro Thr
    370                 375                 380

Phe Gly Asp Leu Leu Arg Glu Glu Tyr Ile Gly Asp Asn Gly Asp Pro
385                 390                 395                 400
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Gln Thr Leu Gln Ala Gln Phe Gln Glu Met Met Ala Asp Ser Met Phe
405 410 415

Val Ile Pro Ala Leu Gln Val Ala His Phe Gln Cys Ser Arg Ala Pro
420 425 430

Val Tyr Phe Tyr Glu Phe Gln His Gln Pro Ser Trp Leu Lys Asn Ile
435 440 445

Arg Pro Pro His Met Lys Ala Asp His Gly Asp Glu Leu Pro Phe Val
450 455 460

Phe Arg Ser Phe Phe Gly Gly Asn Tyr Ile Lys Phe Thr Glu Glu Glu
465 470 475 480

Glu Gln Leu Ser Arg Lys Met Met Lys Tyr Trp Ala Asn Phe Ala Arg
485 490 495

Asn Gly Asn Pro Asn Gly Glu Gly Leu Pro His Trp Pro Leu Phe Asp
500 505 510

Gln Glu Glu Gln Tyr Leu Gln Leu Asn Leu Gln Pro Ala Val Gly Arg
515 520 525

Ala Leu Lys Ala His Arg Leu Gln Phe Trp Lys Lys Ala Leu Pro Gln
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Lys Ile Gln Glu Leu Glu Glu Pro Glu Glu Arg His Thr Glu Leu
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